

OM protein - protein search, using sw model

Run on: April 14, 2006, 20:39:58 ; Search time 190 Seconds
(without alignments)
1244.136 Million cell updates/sec

Title: US-10-796-669A-3
Perfect score: 2769
Sequence: 1 MTKKPARKILSFSTTMRNPK.....FIKKVENSIEWLQINDHLKA 538

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 2443163 seqs, 439378781 residues

Total number of hits satisfying chosen parameters: 2443163

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : A_Geneseq_21:*
1: geneseqp1980s:*
2: geneseqp1990s:*
3: geneseqp2000s:*
4: geneseqp2001s:*
5: geneseqp2002s:*
6: geneseqp2003as:*
7: geneseqp2003bs:*
8: geneseqp2004s:*
9: geneseqp2005s:*

GenCore version 5.1.7
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OM protein - protein search, using sw model

Run on: April 14, 2006, 20:43:48 ; Search time 43 Seconds
(without alignments)
1203.828 Million cell updates/sec

Title: US-10-796-669A-3
Perfect score: 2769
Sequence: 1 MTKKPARKILSFSTTMRNPK.....FIKKVENSIEWLQINDHLKA 538

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : PIR_80:*
1: pirl:*

2: pir2:*
3: pir3:*
4: pir4:*

GenCore version 5.1.7
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OM protein - protein search, using sw model

Run on: April 14, 2006, 20:40:28 ; Search time 235 Seconds
(without alignments)
1615.209 Million cell updates/sec

Title: US-10-796-669A-3
Perfect score: 2769
Sequence: 1 MTKKPKARKILSFSTTMRNPK.....FIKKVENSIEWLQINDHLKA 538

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 2166443 seqs, 705528306 residues

Total number of hits satisfying chosen parameters: 2166443

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : UniProt_05.80:*
1: uniprot_sprot:*
2: uniprot_trembl:*